

SEQUENCE LISTING

19

15

<110>	Glaxo Gro	up Li	mited
	Daniel K.	Burn	s
	Michael P	hilip	Weiner

<120> Multiple Sequencing Method

 \mathcal{B}

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<130> PU3562
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- <140> PCT/US99/21092
- <141> 1999-09-14
- <150> 60100491
- <151> 1998-09-16
- <150> 60100704
- <151> 1998-09-17
- <160> 4
- <170> FastSEQ for Windows Version 4.0
- <210> 1
- <211> 19
- <212> DNA
- <213> Artificial Sequence
- <220>
- <223> homo sapien
- <400> 1
- aattcggctc gagctggag
- <210> 2
- <211> 15
- <212> DNA
- <213> Artificial Sequence
- <220>
- <223> homo sapien
- <400> 2
- ctccagctcg agccg
- <210> 3
- <211> 63
- <212> DNA
- <213> Artificial Sequence
- <220>

<223> DNA fragment not treated with Bpm-I illustrating
sequencing method

<400> 3
gaattcggct cgagctggag actaagttga gatgatatca tttacggggg aaggcgcttt 60
gtg 63

<210> 4
<211> 51
<212> DNA
<213> Artificial Sequence

<220>
<223> DNA fragment treated with Bpm-I illustrating
sequencing method

<221> misc_feature
<222> 35, 36, 39, 40, 43, 44, 47, 49, 50, 51
<223> n = A,T,C or G

gaattcggct cgagctggag gtcaacccaa cacannttnn ttnnggntnn n

51

<400> 4